

(<del>A</del>

CDR2 DASTRAP K,,SLES KV,N,DS K,,SLES ,,,,LES ,,,,LES		CDR2 SRINSDGSSTNYADSVEG GV, IPIRGTA,,, QKFQ, ,V, VPIVGT, KH, QKFQ, AL, KK,,, EKY,, E,, K, GYAH, RV, -AY, NP, LKS	FR4 WGQGTLVTVSS ''',A'',' ''',K,I,L
FR2 WYQQKPGQAPRLIMY ''','','','',',',',' 'I,',R',',P',','I,' ''','','','','','','','','','','','',	FR4  T. FGQGTKLEIKRT P	FR2 WVRQAPGKGLEWV SRIN ''''''''''''''''''''''''''''''''''''	CDR3 SRGGLWDWSPRRIEETKTPFDY WGATG,GRYPTGS,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
CDR1  RAGQSLDSSLLS  ,, SED, NKW, A ,, S, GIS, W, A ,, S, DISIR, N ,, S, GISIR, N	CDR3  AVYY CQQHYNLPRT ,T,,,,YQSY,Y, GL,,,,Y,GVQF,I, ,T,,,,YGSY,L, ,T,,,,YGSY,L,	CDR1 SYWMH ,,GIS RNPIS ,,,,, DFYWS	
FR1ELTQGPATLSLSPGERATLSC RA ELOM,,S,S,,,A,V,D,V,VT,,,,,,S,LS,PVTL,QP,SI,,,S,,,S,S,,A,V,D,V,IT,,, ELOM,,S,SS,,A,V,D,V,IT,,,,,,,S,SS,,A,V,D,V,IT,,,,	FR3  GVPARFSGSGSGTDFTLTISSLQPEDFAVYY  ,,,S,,,,,,A,,,,K,TRVEA,,VGL,,  ,,S,,,,,,B,,,,,,T,,,T,,T,,T,,T,,T,,T,,T,,T,	FR1  EVQLLES-GGGLVQPGGSLRLSCAASGVTFS  ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	FR3  RFTISRDNAKNTLYLQMNSLRAEDTAVYYC ,V,YTA,ESTS,V,MELS,,,S,,,,,,,,,,,,,,,S,,,,,S,,,,,,,,,,
5A7 3C1. 3E4 7G4 5H2 5D9	5A7 3C1 3B4 7G4 5H2 5D9	(B) 5A7 3C1 3E4 7G4 5H2 5D9	5A7 3C1 3E4 7G4 5H2 5D9

Fig. 2



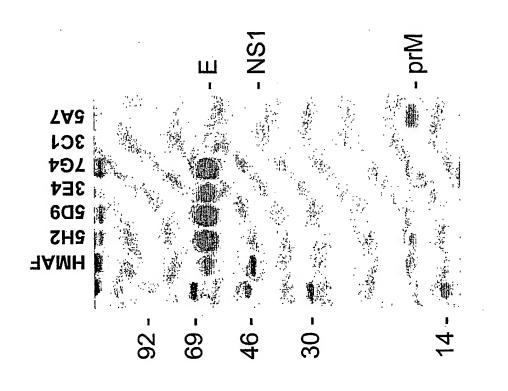
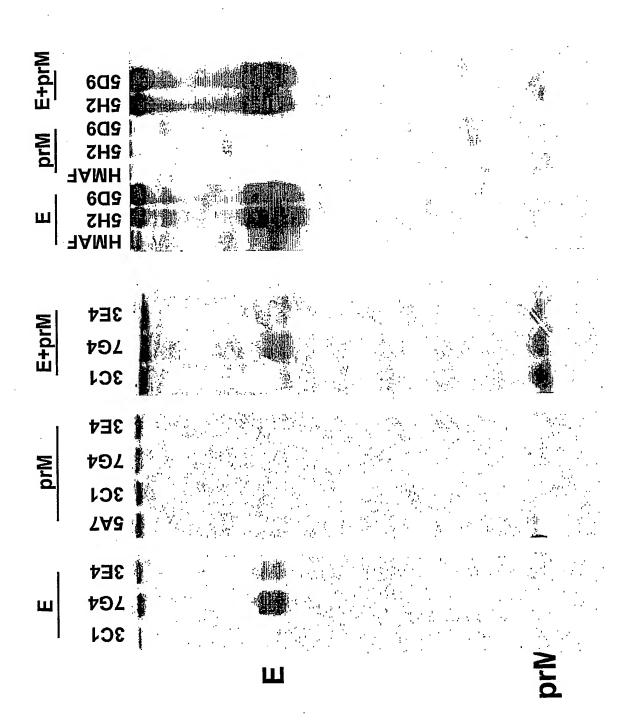
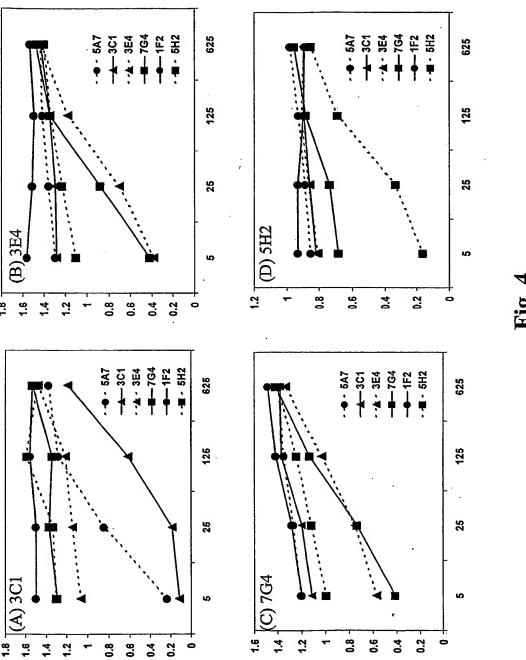


Fig. 3A





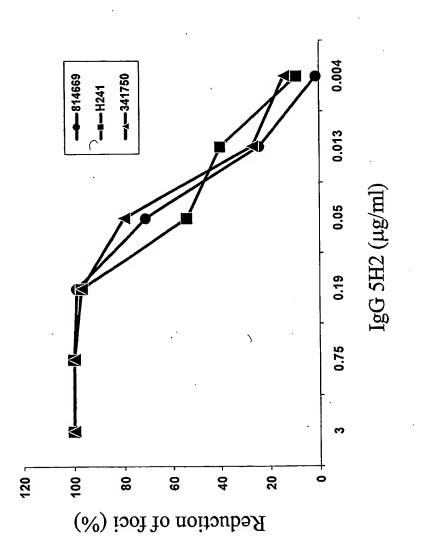
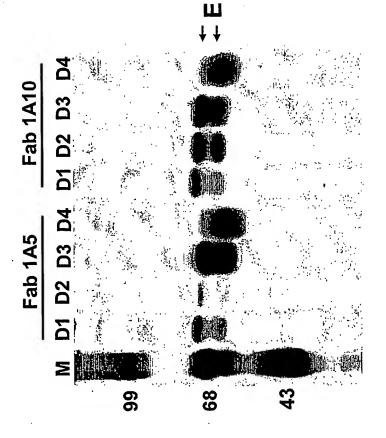


Fig. 5

V. M. VTAA	VV.IIA.ESTSTAYMELSSSTYYADVSSYSEY	ASV.AGMPAAGIL SEISTAYMELSSSTYYADSSYSE.
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CDR2 VSSTLQS FAH. HA HA DA.S.E. GLSNRA.		CDR2	ALIYSAD-STHYADSVKGT GY.SYRG-T.Y.NP.L.S GV.VPIVGT.KH.QKFQ.	88
FR2 WYQQKPGKAPKLLIS	FR4  9QLEV QLE QLE  WQLE  WQLE  YQLE	FR2	DNVMH WVRQAPGKGLEWV ALIYSAD-STHYADSVKG	CDR3 FR4  AREYCTGGT-CFAHFDY WGQGTLUVTVSS
CDR1  RASQSITNYLS TV. G.SSE.N  KSLLHSDGNT.F	CDR3 LTISSLQPEDFATYY CHYG-YGTHTNDQQDQHFNSFPWX.QVEAVGVFMQ.TQLPYX.TRVEAVGLYVQ.VQFPI.	CDR1		CDR3  DTAVYYC AREYCTGGT-CFAHFDY
FR1  ELQMTQSPSSLSASVGDRVTITCELVTA. ELVAP.QPAS.S.	FR3  GVP SRFSGSGSGTDFTLTISSLQPEDFATYY  T	FR1	EVQLLE-SGGGLVQPGGSRRLSCAASGFTISQ	FR3   CDR3
(A) 2H7 2H5 3A2 1A5 1B2 1A10 3E4	2H7 2H5 3A2 1A5 1B2 1A10 3B4	(B)	2H7 2H5 3A2 1A5 1B2 1A10	2H7 2H5 3A2 1A5 1B2 1A10





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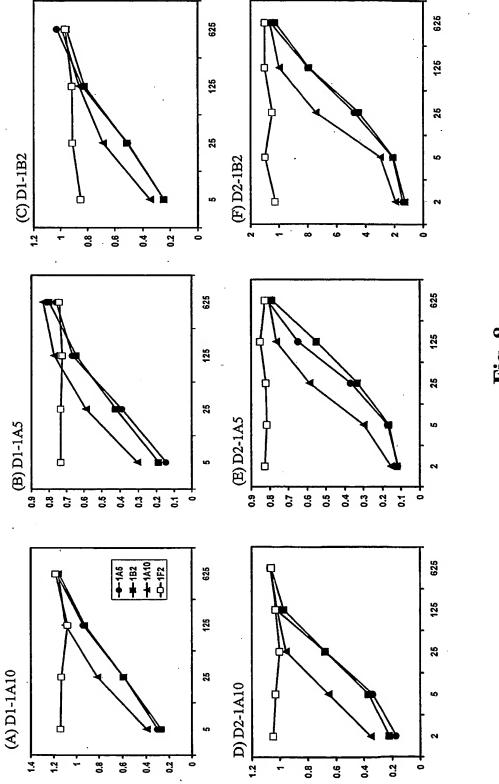


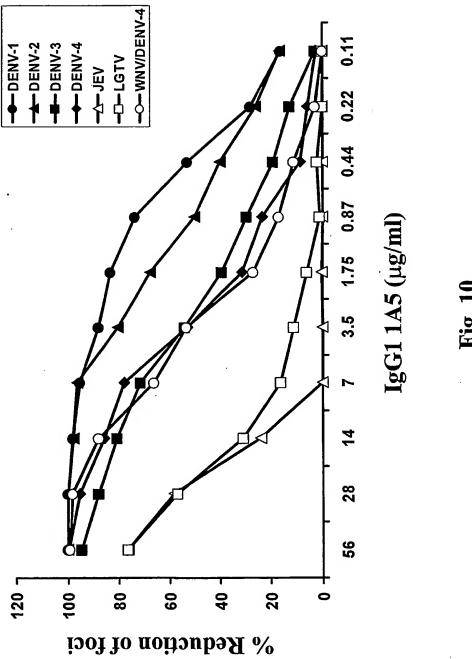
Fig. 8



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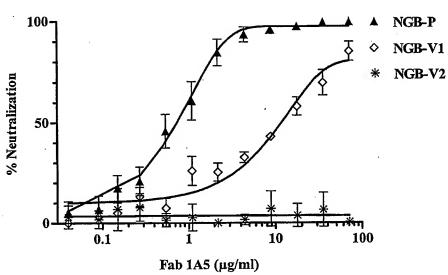


Fig. 5

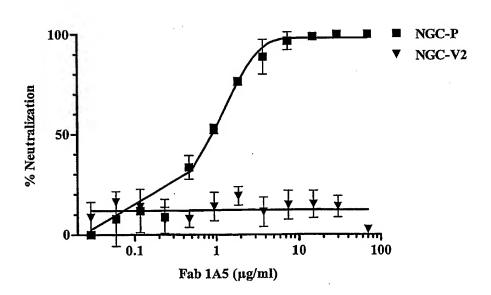


.

A



В



**Fig. 11** 

A

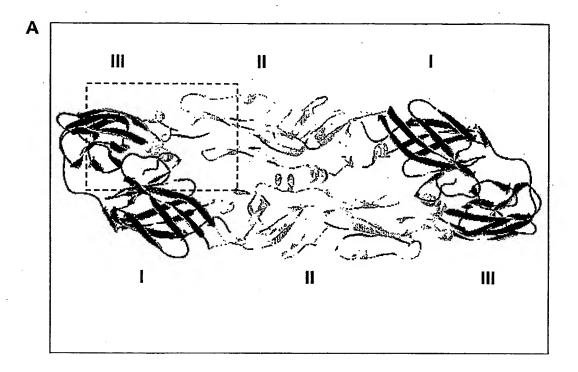
			$Gly_{106}$		
		C	$\downarrow$	d	
DENV-2 P	89	REVCKHSMVDRG	WGNGCGLFGK	GGIVTCAMFT	120
DENV-2 V1					
DENV-2 V2			Ÿ		
DENV-1		NRRTF		-SLIK-K	
DENV-3		NYTY		-SLK-Q	
DENV-4		QYI-RRDV		VK-S	
WNV		ARQGV		-S-DK-A	
JEV		SYQGFT		-S-DK-S	
JEV SA14-14-2		SYQGFT	F	-S-DK-S	
SLEV		TRDV		-s-DK	
YFV Asibi		DNARTYS		-SAK	
YFV 17D		DNARTYS		-SAK	
LGTV		GTRDQS	H	-SVK	
TBEV		GTRDQS	H	-SA-VKAA	

 $\mathbf{B}$ 

	His <sub>317</sub>
	$\mathbf{A}$ $\mathbf{b}$ $\mathbf{B}$
<b>DENV-2</b> P 306	FKVVKEIAETQHGTIVIRVQY 326
DENV-2 V1	Q
DENV-2 V2	
DENV-1	LEVVLVQ-K-
DENV-3	-VLKVSL-K-E-
DENV-4	-SIDMT-VK-K-
WNV	FLGTP-D-GV-LEL
JEV	-SFA-NP-D-GVELT-
JEV SA14-14-2	-SFA-NP-D-GVELT-
SLEV	-TFS-NPTD-GVIVEL
YFV Asibi	MFFNPDT-GV-MQ-KV
YFV 17D	MFFNPDT-GV-MQ-KV
LGTV	-TWKRAPTDSG-D-V-ME-GF
TREV	-TWRRAPTDSG-D-V-ME-TF

Fig. 12





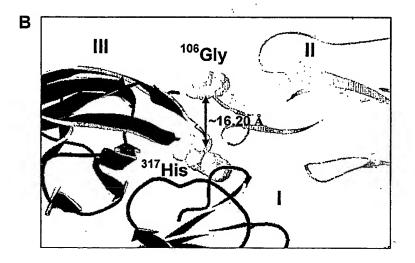
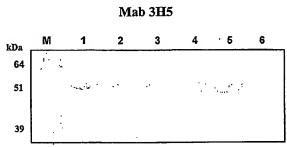
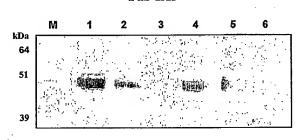


Fig. 13

A



Fab 1A5



B

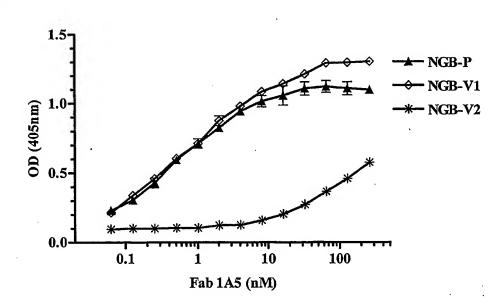


FIG. 14

Fig. 14

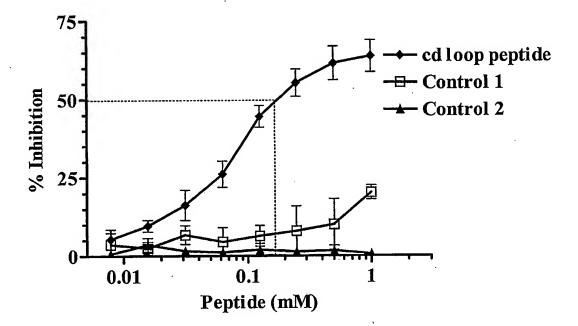
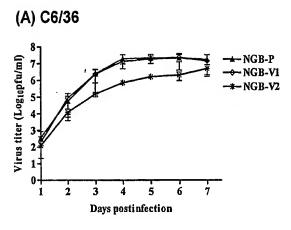
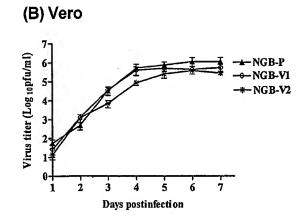
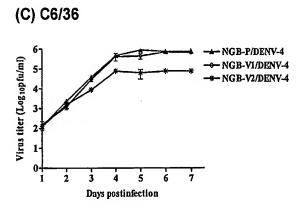
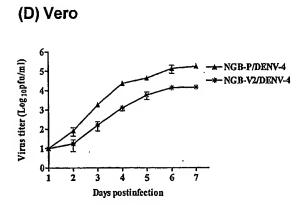


Fig. 15

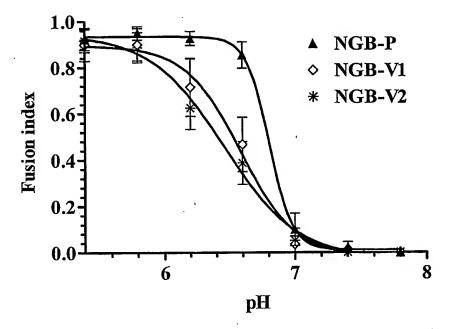




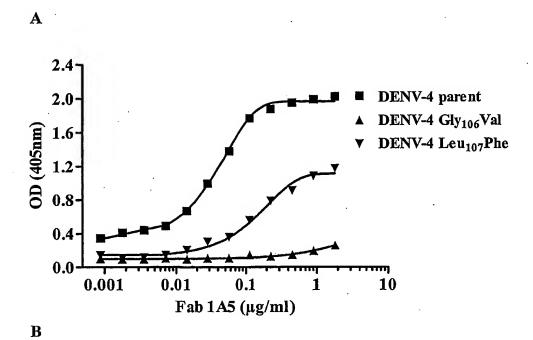




**Fig. 16** 



**Fig. 17** 



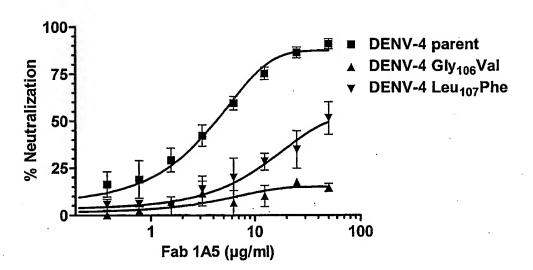


Fig. 18